	CRF Errors Corrocted by the STIC Systems Branch  CRF Processing Date: 10/23/20  Tumber: 0.9/936,677
באפור ע	Changed a lile from non-ASCII to ASCII ENTERE WHOO by: 100 STIC
	Changed the margins in cases where the sequence text was "wrapped" down to the next line.
].	Edited a format error in the Current Application Data section, specifically:
) <i>.</i> ,	Edited the Current Application Data section with the actual current number. The number inputted by the applicant was   The prior application data; or other
	Added the mandatory heading and subheadings for "Current Application Data".
	Edited the 'Number of Sequences' field. The applicant spelled out a number instead of using an integer.
	Changed the spelling of a mandatory field (the headings or subheadings), specifically:
	Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were edited were:
	Inserted or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited:
(	Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
	Inserted colons after headings/subheadings. Headings edited included: •, . * .
-	Deleted extra, invalid, headings-used by an applicant, specifically:
	Deleted: non-ASCII garbage at the beginning/end of files: secretary initials/filename at end of file page numbers throughout text: other invalid text, such as
	Inserted mandatory headings, specifically:
(	Corrected an obvious erro: in the response, specifically:
	Edited identifiers where upper case is used but lower case is required, or vice versa.
(	Corrected an error in the Number of Sequences field, specifically:
_	Mand Pago Break gode was inserted by the applicant. All occurrences had to be deleted.
	eloted ending stop codon in amino acid sequences and adjusted the "(A)Length: field accordingly (error o to a Patentin bug). Sequences corrected:
	Other:
 :	
	OWIGO

Examiner: The above corrections must be communicated to the applicant in the first Office Action! DO NOT send a copy of this form.

Input Set : A:\PTO.AMC.txt

```
3 <110> APPLICANT: LIBON Christine
             CORVAIA Nathalie
             BECK Alain
             BONNEFOY Jean-Yves
       <120> TITLE OF INVENTION: IMMUNOSTIMULATING BACTERIAL MEMBRANE FRACTIONS IN CANCER
              TREATMENT
    11 <130> FILE REFERENCE: D17974
C--> 13 <140> CURRENT APPLICATION NUMBER: US/09/936,677
C--> 13 <141> CURRENT FILING DATE: 2001-09-14
    13 <150> PRIOR APPLICATION NUMBER: FR 99 03 154
    14 <151> PRIOR FILING DATE: 1999-03-15
    16 <150> PRIOR APPLICATION NUMBER: PCT/FR00/00623
    17 <151> PRIOR FILING DATE: 2000-03-15
    19 <160> NUMBER OF SEQ ID NOS: 4
     21 <170> SOFTWARE: PatentIn Vers. 2.0
     23 <210> SEO ID NO: 1
    24 <211> LENGTH: 1035
    25 <212> TYPE: DNA
    26 <213> ORGANISM: Klebsiella pneumoniae
     28 <220> FEATURE:
     29 <221> NAME/KEY: exon
     30 <222> LOCATION: (1)..(1032)
     32 <220> FEATURE:
     33 <221> NAME/KEY: intron
     34 <222> LOCATION: (1033)..(1035)
     36 <220> FEATURE:
     37 <221> NAME/KEY: CDS
     38 <222> LOCATION: (1)..(1032)
     40 <400> SEQUENCE: 1
     41 atg aaa gca att ttc gta ctg aat gcg gct ccg aaa gat aac acc tgg
    42 Met Lys Ala Ile Phe Val Leu Asn Ala Ala Pro Lys Asp Asn Thr Trp
     43
                          5
         1
     45 tat gca ggt ggt aaa ctg ggt tgg tcc cag tat cac gac acc ggt ttc
     46 Tyr Ala Gly Gly Lys Leu Gly Trp Ser Gln Tyr His Asp Thr Gly Phe
                                         25
     49 tac ggt aac ggt ttc cag aac aac ggt ccg acc cgt aac gat cag
     50 Tyr Gly Asn Gly Phe Gln Asn Asn Gly Pro Thr Arg Asn Asp Gln
                                     40
     53 ctt ggt gct ggt gcg ttc ggt ggt tac cag gtt aac ccg tac ctc ggt
     54 Leu Gly Ala Gly Ala Phe Gly Gly Tyr Gln Val Asn Pro Tyr Leu Gly
             50
                                 55
     57 ttc gaa atg ggt tat gac tgg ctg ggc cgt atg gca tat aaa ggc agc
     58 Phe Glu Met Gly Tyr Asp Trp Leu Gly Arg Met Ala Tyr Lys Gly Ser
                                                 75
                                                                           288
     61 gtt gac aac ggt gct ttc aaa gct cag ggc gtt cag ctg acc gct aaa
     62 Val Asp Asn Gly Ala Phe Lys Ala Gln Gly Val Gln Leu Thr Ala Lys
                         85
```

Input Set : A:\PTO.AMC.txt

65 ctg ggt tac 66 Leu Gly Tyr 67			Asp Ile Tyr				
69 ggc atg gtt 70 Gly Met Val 71 115	tgg cgc gct	gac tcc aaa	ggc aac tac				
7.3 gtt tcc cgt 74 Val Ser Arg 75 130				Val Phe Ala			
77 ggc gta gag 78 Gly Val Glu 79 145							
81 cag tgg gtt 82 Gln Trp Val 83							
85 gat aac ggc 86 Asp Asn Gly 87	Met Leu Ser 180	Leu Gly Val	Ser Tyr Arg	y Phe Gly Gln 190	Glu		
89 gat gct gca 90 Asp Ala Ala 91 195	Pro Val Val	Ala Pro Ala 200	Pro Ala Pro	Ala Pro Glu 205	Val		
93 gct acc aag 94 Ala Thr Lys 95 210	His Phe Thr	Leu Lys Ser 215	Asp Val Leu 220	Phe Asn Phe	Asn		
97 aaa gct acc 98 Lys Ala Thr 99 225	Leu Lys Pro 230	Glu Gly Glr	Gln Ala Leu 235	ı Asp Gln Leu	Tyr 240		
101 act cag cto 102 Thr Gln Let 103	ı Ser Asn Me 245	t Asp Pro Ly	s Asp Gly Se 250	er Ala Val Val 25	l Leu 5		
105 ggc tac acc 106 Gly Tyr Thi 107	r Asp Arg Ilo 260	e Gly Ser Gl 26	u Ala Tyr As 55	sn Gln Gln Le 270	u Ser		
109 gag aaa cgt 110 Glu Lys Arg 111 275	g Ala Gln Se: 5	r Val Val As 280	p Tyr Leu Va	al Ala Lys Gly 285	y Ile		
113 ccg gct ggc 114 Pro Ala Gly 115 290	y Lys Ile Se	r Ala Arg Gl 295	y Met Gly Gl. 30	Lu Ser Asn Pro 00	o Val		
117 act ggc aad 118 Thr Gly Asi 119 305	n Thr Cys Asy	p Asn Val Ly O	s Ala Arg Al 315	la Ala Leu Ilo	e Asp 320		
121 tgc ctg gct 122 Cys Leu Ala -123	t ccg gat cg a Pro Asp Ar 325	t cgt gta ga g Arg Val Gl	ng atc gaa gt Lu Ile Glu Va 330	t aaa ggc ta al Lys Gly Ty: 33	r Lys		
125 gaa gtt gta 126 Glu Val Val 127			ıa		1035		
130 <210> SEQ ID NO: 2							

Input Set : A:\PTO.AMC.txt

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131 <211> LENGTH: 344
132 <212> TYPE: PRT
133 <213> ORGANISM: Klebsiella pneumoniae
135 <400> SEQUENCE: 2
136 Met Lys Ala Ile Phe Val Leu Asn Ala Ala Pro Lys Asp Asn Thr Trp
139 Tyr Ala Gly Gly Lys Leu Gly Trp Ser Gln Tyr His Asp Thr Gly Phe
                20
142 Tyr Gly Asn Gly Phe Gln Asn Asn Gly Pro Thr Arg Asn Asp Gln
143 35
                                 40
145 Leu Gly Ala Gly Ala Phe Gly Gly Tyr Gln Val Asn Pro Tyr Leu Gly
148 Phe Glu Met Gly Tyr Asp Trp Leu Gly Arg Met Ala Tyr Lys Gly Ser
                         70
151 Val Asp Asn Gly Ala Phe Lys Ala Gln Gly Val Gln Leu Thr Ala Lys
154 Leu Gly Tyr Pro Ile Thr Asp Asp Leu Asp Ile Tyr Thr Arg Leu Gly
                100
                                   105
157 Gly Met Val Trp Arg Ala Asp Ser Lys Gly Asn Tyr Ala Ser Thr Gly
           115
                               120
160 Val Ser Arg Ser Glu His Asp Thr Gly Val Ser Pro Val Phe Ala Gly
                            135
163 Gly Val Glu Trp Ala Val Thr Arg Asp Ile Ala Thr Arg Leu Glu Tyr
                        150
                                            155
166 Gln Trp Val Asn Asn Ile Gly Asp Ala Gly Thr Val Gly Thr Arg Pro
                                        170
                    165
169 Asp Asn Gly Met Leu Ser Leu Gly Val Ser Tyr Arg Phe Gly Gln Glu
               180
                                    185
172 Asp Ala Ala Pro Val Val Ala Pro Ala Pro Ala Pro Ala Pro Glu Val
           195
                                200
175 Ala Thr Lys His Phe Thr Leu Lys Ser Asp Val Leu Phe Asn Phe Asn
176 210
                            215
178 Lys Ala Thr Leu Lys Pro Glu Gly Gln Gln Ala Leu Asp Gln Leu Tyr
                                            235
181 Thr Gln Leu Ser Asn Met Asp Pro Lys Asp Gly Ser Ala Val Val Leu
184 Gly Tyr Thr Asp Arg Ile Gly Ser Glu Ala Tyr Asn Gln Gln Leu Ser
                                    265
                260
187 Glu Lys Arg Ala Gln Ser Val Val Asp Tyr Leu Val Ala Lys Gly Ile
                                280
190 Pro Ala Gly Lys Ile Ser Ala Arg Gly Met Gly Glu Ser Asn Pro Val
                            295
                                                300
        290
193 Thr Gly Asn Thr Cys Asp Asn Val Lys Ala Arg Ala Ala Leu Ile Asp
                        310
                                            315
196 Cys Leu Ala Pro Asp Arg Arg Val Glu Ile Glu Val Lys Gly Tyr Lys
                                        330
                    325
199 Glu Val Val Thr Gln Pro Ala Gly
                340
 203 <210> SEQ ID NO: 3
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Input Set : A:\PTO.AMC.txt

```
204 <211> LENGTH: 303
 205 <212> TYPE: DNA
 206 <213> ORGANISM: Klebsiella pneumoniae
 208 <220> FEATURE:
 209 <221> NAME/KEY: CDS
 210 <222> LOCATION: (1)..(303)
212 <400> SEQUENCE: 3
 213 acc gtg aaa acc aaa aac acc acg acc cag acc cag ccg agc aaa
                                                                  48
 214 Thr Val Lys Thr Lys Asn Thr Thr Thr Gln Thr Gln Pro Ser Lys
 215
                   5
 96
 218 Pro Thr Thr Lys Gln Arg Gln Asn Lys Pro Pro Asn Lys Pro Asn Asn
                                   25
 221 gat ttc cat ttc gaa gtg ttc aac ttc gtg ccg tgc agc atc tgc agc
                                                                  144
 222 Asp Phe His Phe Glu Val Phe Asn Phe Val Pro Cys Ser Ile Cys Ser
                               40
 225 aac aac ccg acc tgc tgg gcg atc tgc aaa cgt atc ccg aac aaa aaa
                                                                  192
 226 Asn Asn Pro Thr Cys Trp Ala Ile Cys Lys Arg Ile Pro Asn Lys Lys
         50
                           55
 230 Pro Gly Lys Lys Thr Thr Lys Pro Thr Lys Lys Pro Thr Phe Lys
                        70
                                          75
 233 acc acc aaa aaa gat cat aaa ccg cag acc acc aaa ccg aaa gaa gtg
 234 Thr Thr Lys Lys Asp His Lys Pro Gln Thr Thr Lys Pro Lys Glu Val
 235
                    85
                                      90
                                                                  303
 237 ccg acc acc aaa ccg
 238 Pro Thr Thr Lys Pro
 239
               100
 242 <210> SEQ ID NO: 4
 243 <211> LENGTH: 101
 244 <212> TYPE: PRT
 245 <213> ORGANISM: Klebsiella pneumoniae
247 <400> SEQUENCE: 4
 248 Thr Val Lys Thr Lys Asn Thr Thr Thr Thr Gln Thr Gln Pro Ser Lys
                                      10
 251 Pro Thr Thr Lys Gln Arg Gln Asn Lys Pro Pro Asn Lys Pro Asn Asn
                20
                                   25
 254 Asp Phe His Phe Glu Val Phe Asn Phe Val Pro Cys Ser Ile Cys Ser
             35
                               40
 257 Asn Asn Pro Thr Cys Trp Ala Ile Cys Lys Arg Ile Pro Asn Lys Lys
                           55
 260 Pro Gly Lys Lys Thr Thr Lys Pro Thr Lys Lys Pro Thr Phe Lys
                       70
                                          75
 263 Thr Thr Lys Lys Asp His Lys Pro Gln Thr Thr Lys Pro Lys Glu Val
                                      90
 266 Pro Thr Thr Lys Pro
 267
```

VERIFICATION SUMMARY DATE: 10/23/2001 PATENT APPLICATION: US/09/936,677 TIME: 13:31:58

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\10232001\1936677.raw

L:13 M:270 C: Current Application Number differs, Replaced Current Application No

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date